

INPUT SET: S7400.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/444,994DATE: 11 21 95
TIME: 08:56:58

INPUT SET: S7400.raw

47 (A) LENGTH: 19 base pairs
48 (B) TYPE: nucleic acid
49 (C) STRANDEDNESS: single
50 (D) TOPOLOGY: linear
51
52 (ii) MOLECULE TYPE: DNA
53
54
55
56
57 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
58
59 GCAAAGCAGG AGAAACCAC 19
60
61 (2) INFORMATION FOR SEQ ID NO:2:
62
63 (i) SEQUENCE CHARACTERISTICS:
64 (A) LENGTH: 24 base pairs
65 (B) TYPE: nucleic acid
66 (C) STRANDEDNESS: single
67 (D) TOPOLOGY: linear
68
69 (ii) MOLECULE TYPE: DNA
70
71
72
73
74 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
75
76 GGGTCCATCT GATAGATATG AGAG 24
77
78 (2) INFORMATION FOR SEQ ID NO:3:
79
80 (i) SEQUENCE CHARACTERISTICS:
81 (A) LENGTH: 48 base pairs
82 (B) TYPE: nucleic acid
83 (C) STRANDEDNESS: single
84 (D) TOPOLOGY: linear
85
86 (ii) MOLECULE TYPE: DNA
87
88
89 (ix) FEATURE:
90 (A) NAME/KEY: modified_base
91 (B) LOCATION: 36
92 (D) OTHER INFORMATION: /mod_base= i
93
94 (ix) FEATURE:
95 (A) NAME/KEY: modified_base
96 (B) LOCATION: 37
97 (D) OTHER INFORMATION: /mod_base= i
98
99 (ix) FEATURE:

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/444,994DATE: 11 21 95
TIME: 08:57:02

INPUT SET: S7400.raw

100 (A) NAME/KEY: modified_base
101 (B) LOCATION: 41
102 (D) OTHER INFORMATION: /mod_base= i
103
104 (ix) FEATURE:
105 (A) NAME/KEY: modified_base
106 (B) LOCATION: 42
107 (D) OTHER INFORMATION: /mod_base= i
108
109 (ix) FEATURE:
110 (A) NAME/KEY: modified_base
111 (B) LOCATION: 46
112 (D) OTHER INFORMATION: /mod_base= i
113
114 (ix) FEATURE:
115 (A) NAME/KEY: modified_base
116 (B) LOCATION: 47
117 (D) OTHER INFORMATION: /mod_base= i
118
119
120 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
121
122 CUACUACUAC UAGGCCACGC GTCCACTACT ACGGNNNGGG NNGGGNNG 48
123
124 (2) INFORMATION FOR SEQ ID NO:4:
125
126 (i) SEQUENCE CHARACTERISTICS:
127 (A) LENGTH: 20 base pairs
128 (B) TYPE: nucleic acid
129 (C) STRANDEDNESS: single
130 (D) TOPOLOGY: linear
131
132 (ii) MOLECULE TYPE: DNA
133
134
135
136
137 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
138
139 TCCTGATGTT GCTGTAGACG 20
140
141 (2) INFORMATION FOR SEQ ID NO:5:
142
143 (i) SEQUENCE CHARACTERISTICS:
144 (A) LENGTH: 20 base pairs
145 (B) TYPE: nucleic acid
146 (C) STRANDEDNESS: single
147 (D) TOPOLOGY: linear
148
149 (ii) MOLECULE TYPE: DNA
150
151
152

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/444,994DATE: 11/21/95
TIME: 08:57:06

INPUT SET: S7400.raw

153
154 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
155
156 GCACGACTAG TATGATTTGC 20
157
158 (2) INFORMATION FOR SEQ ID NO:6:
159
160 (i) SEQUENCE CHARACTERISTICS:
161 (A) LENGTH: 8 amino acids
162 (B) TYPE: amino acid
163 (C) STRANDEDNESS:
164 (D) TOPOLOGY: unknown
165
166 (ii) MOLECULE TYPE: peptide
167
168
169
170
171 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
172
173 Thr Gly Ala Gly Ala Gly Leu Gly
174 1 5
175
176 (2) INFORMATION FOR SEQ ID NO:7:
177
178 (i) SEQUENCE CHARACTERISTICS:
179 (A) LENGTH: 5 amino acids
180 (B) TYPE: amino acid
181 (C) STRANDEDNESS:
182 (D) TOPOLOGY: unknown
183
184 (ii) MOLECULE TYPE: peptide
185
186
187
188
189 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
190
191 Tyr Ser Ala Ala Lys
192 1 5
193
194 (2) INFORMATION FOR SEQ ID NO:8:
195
196 (i) SEQUENCE CHARACTERISTICS:
197 (A) LENGTH: 27 base pairs
198 (B) TYPE: nucleic acid
199 (C) STRANDEDNESS: single
200 (D) TOPOLOGY: unknown
201
202 (ii) MOLECULE TYPE: DNA
203
204
205 (ix) FEATURE:

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/444,994DATE: 11/21/95
TIME: 08:57:09

INPUT SET: S7400.raw

206 (A) NAME/KEY: CDS
207 (B) LOCATION: 1..27
208
209
210 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8: 27
211
212 GAC TGG CTG GAA TTC CCC ATG GCG TCC
213 Asp Trp Leu Glu Phe Pro Met Ala Ser
214 1 5
215
216
217 (2) INFORMATION FOR SEQ ID NO:9:
218
219 (i) SEQUENCE CHARACTERISTICS:
220 (A) LENGTH: 9 amino acids
221 (B) TYPE: amino acid
222 (D) TOPOLOGY: linear
223
224 (ii) MOLECULE TYPE: protein
225
226 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:
227
228 Asp Trp Leu Glu Phe Pro Met Ala Ser
229 1 5
230
231 (2) INFORMATION FOR SEQ ID NO:10:
232
233 (i) SEQUENCE CHARACTERISTICS:
234 (A) LENGTH: 2940 base pairs
235 (B) TYPE: nucleic acid
236 (C) STRANDEDNESS: unknown
237 (D) TOPOLOGY: unknown
238
239 (ii) MOLECULE TYPE: cDNA
240
241
242 (ix) FEATURE:
243 (A) NAME/KEY: CDS
244 (B) LOCATION: 47..1663
245
246
247 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:
248
249 CTAACCTCAG CGGTGGCACC GGGATCGGTT GCCTTGAGCC TGAAAT ATG ACC ACC 55
250 Met Thr Thr
251 1
252
253 CCA GGA AAA GAG AAC TTT CGC CTG AAA AGT TAC AAG AAC AAA TCT CTG 103
254 Pro Gly Lys Glu Asn Phe Arg Leu Lys Ser Tyr Lys Asn Lys Ser Leu
255 5 10 15
256
257 AAT CCC GAT GAG ATG CGC AGG AGG AGG GAG GAA CAA GCA CTG CAG TTA 151
258 Asn Pro Asp Glu Met Arg Arg Glu Glu Gly Leu Gln Leu

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION **US/08/444,994**

DATE: 11/21/95
TIME: 08:57:12

INPUT SET: S7400.raw

Line	Error	Original Text
------	-------	---------------

542	Stop Codon at end of sequence removed - no error	
929	Stop Codon at end of sequence removed - no error	
1376	Stop Codon at end of sequence removed - no error	

NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 CFR 1.821 - 1.825 for the following reason(s):

1. This application clearly fails to comply with the requirements of 37 CFR 1.821 - 1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.

2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 CFR 1.821(c).

3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 CFR 1.821.

4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 CFR 1.822 and/or 1.823, as indicated on the attached marked-up copy of the "Raw Sequence Listing."

5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A substitute computer readable form must be submitted as required by 37 CFR 1.825(d).

6. The paper copy of the "Sequence Listing" is not the same as the computer readable form of the "Sequence Listing" as required by 37 CFR 1.821(e).

7. Other: _____

Applicant must provide:

An initial or substitute computer readable form (CRF) copy of the "Sequence Listing"

An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification

A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 CFR 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d)

For questions regarding compliance with these requirements, please contact:

For Rules Interpretation, call (703) 308-1123

For CRF submission help, call (703) 308-4212

For PatentIn software help, call (703) 308-6856

Please return a copy of this notice with your response.